```
                          FR1                    CDR1              FR2
VHH#3G    QVQLQDSGGGLVQAGGSLRLSCAVSGR  TFSAHS--VYTMG  WFRQAPGKEREFVA
VHH#3E    QVQLQESGGGLVQPGGSLRLSCAASGR  TFSDHSGYTYTIG  WFRQAPGKEREFVA
VHH#1A    QVQLQESGGGLVQPGGSLRLSCATSGF  DFS--VSWMY---  WVRQAPGKGLEWVS
VHH#2B    QVQLQESGGGLVQPGGSLRLSCATSGF  TFS--DYWMY---  WVRQAPGKGLEWVS
VHH#12B   QVQLQESGGGLVQPGGSLRLSCAASGF  EFE--NHWMY---  WVRQAPGKGLEWVS
VHH#7B    QVQLQESGGGLVQPGGSLRLSCAASGS  IFRVNA-----MG  WYRQVPGNQREFVA
          ***** ****** ********* **      *          * ** **    * *


                   CDR2                    FR2
VHH#3G    RIYWSSANTYYADSVKG  RFTISRDNAKNTVDLLMNSLKPEDTAVYYCAA
VHH#3E    RIYWSSGNTYYADSVKG  RFAISRDIAKNTVDLTMNNLEPEDTAVYYCAA
VHH#1A    EINTNGLITKYVDSVKG  RFTISRDNAKNTLYLQMDSLIPEDTALYYCAR
VHH#2B    TVNTNGLITRYADSVKG  RFTISRDNAKYTLYLQMNSLKSEDTAVYYCTK
VHH#12B   TVNTNGLITRYADSVKG  RFTISRDNAKYTLYLQMNSLKSEDTAVYYCTK
VHH#7B    -IITSGDNLNYADAVKG  RFTISTDNVKKTVYLQMNVLKPEDTAVYYCNA
            *  * *** **  **  *    * *    * *   *   **** ***


                CDR3             FR4         Hinge
VHH#3G    RDGIPTSRTVGSYNY  WGQGTQVTVSS  EPKTPKPQP
VHH#3E    RDGIPTSRSVESYNY  WGQGTQVTVSS  EPKTPKPQP
VHH#1A    ----------SPSGSF  RGQGTQVTVSS  EPKTPKPQP
VHH#2B    -VVPPYSDDSRTNAD  WGQGTQVTVSS  EPKTPKPQP
VHH#12B   -VLPPYSDDSRTNAD  WGQGTQVTVSS  EPKTPKPQP
VHH#7B    --ILQTSRWSIPSNY  WGQGTQVTVSS  EPKTPKPQP
                           *********  *********
```
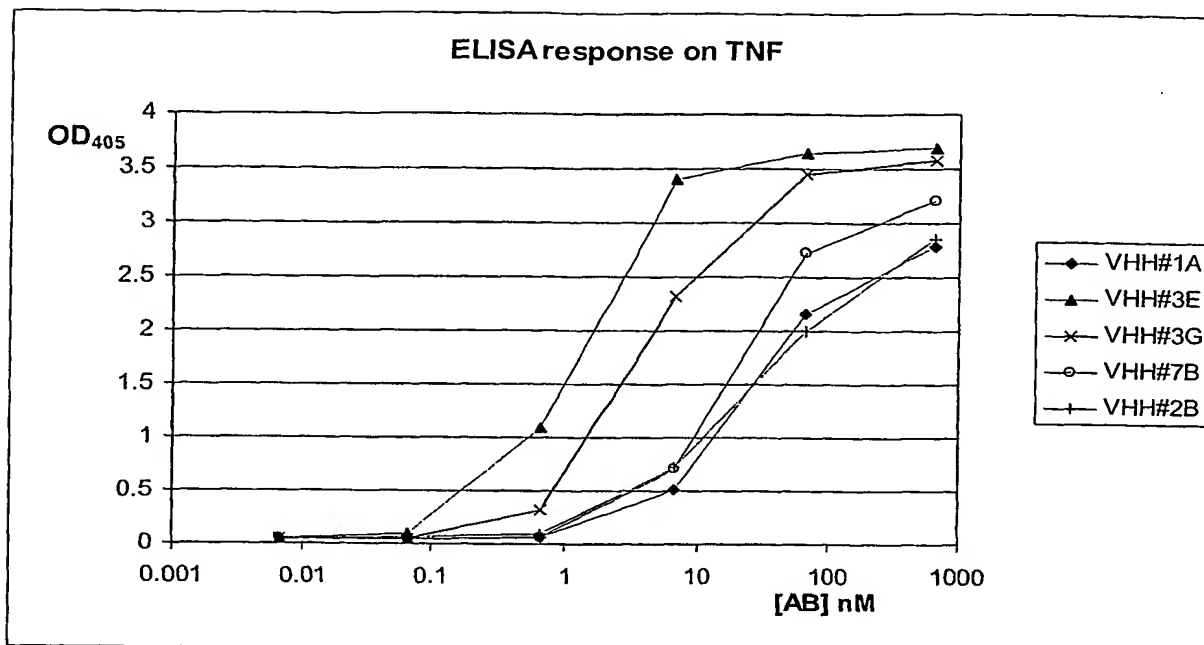
Figure 1

Figure 2

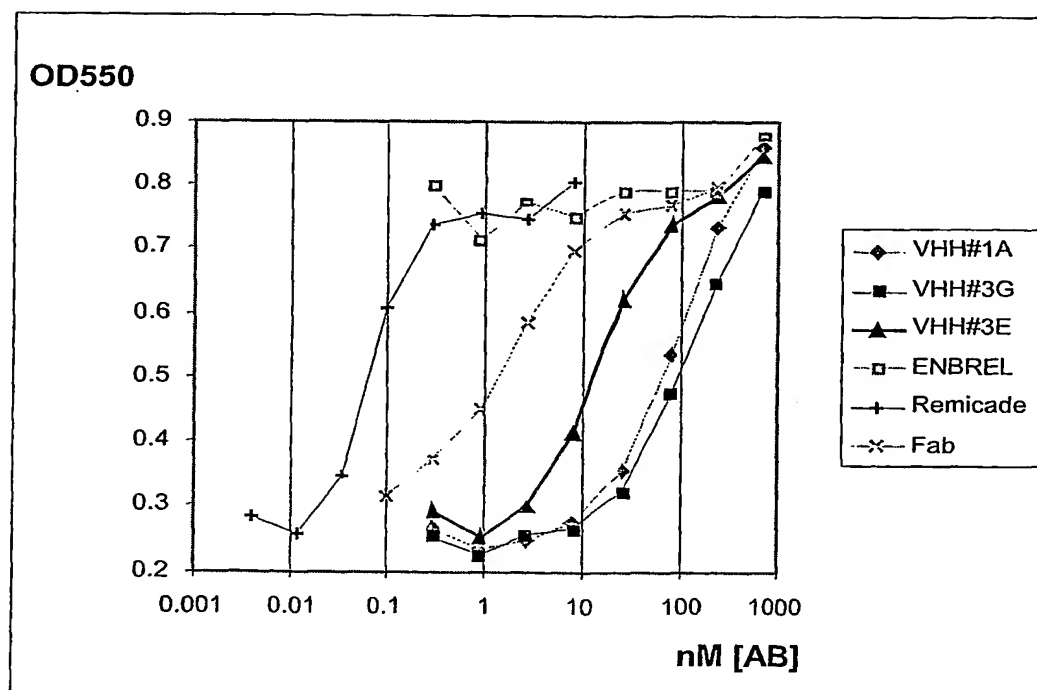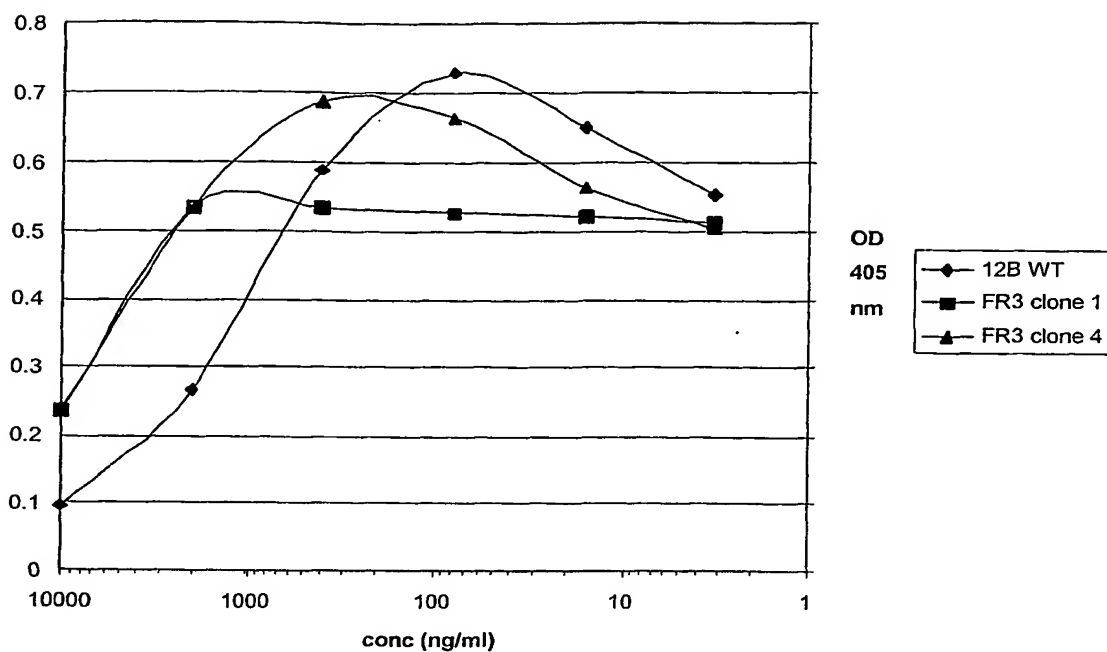**Figure 3**

Figure 4

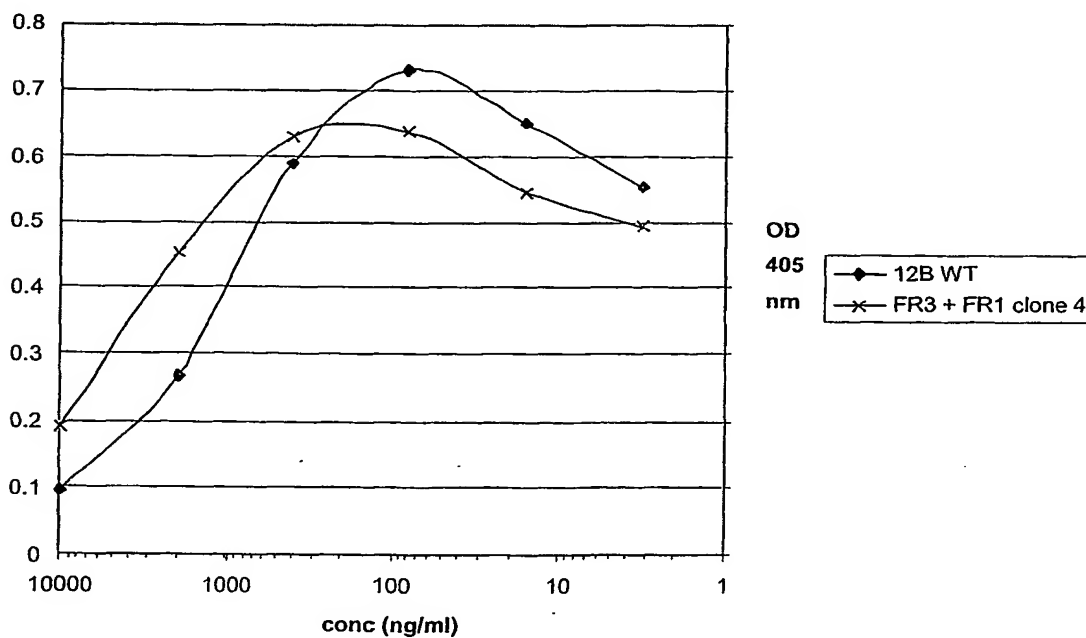Figure 5

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Figure 6

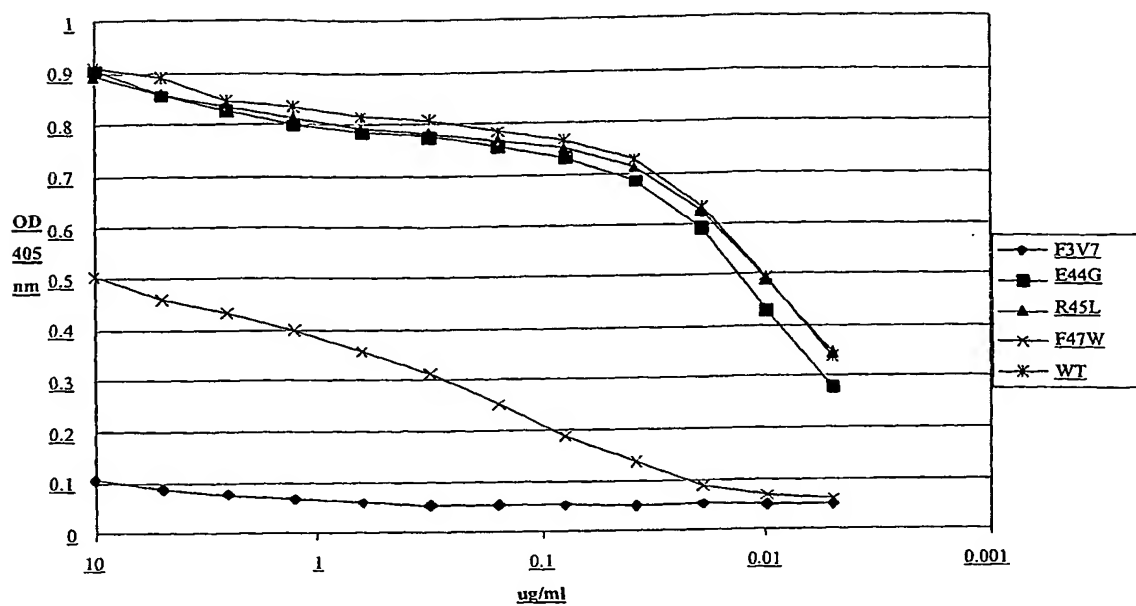Figure 7

```
                            FR1                          CDR1        FR2
VHH#m3F     QVQLQDSGGGLVQAGGSLRLSCAASGG  TFSSIIMA  WFRQAPGKEREFVGA
VHH#m4B     QVQLQDSGGGLVQAGGSLRLSCGVSGL  SFSGYTMG  WFRQAPGKEREFAAA
VHH#m9A     EVQLVESGGGLVQAGGSLRLSCAASGG  TLSSYITG  WFRQAPGKEREFVGA
VHH#m9E     QVQLVESGGGLVQAGGSLRLSCAASEG  TLSGYILG  WFRQAPGKEREFVGA
            ***  *************  *        *         * ***********  *

                      CDR2                             FR2
VHH#m3F     VSWSGGTTVYADSVLG  RFEISRDSARKSVYLQMNSLKPEDTAVYYCAA
VHH#m4B     IGWNSGTTEYRNSVKG  RFTISRDNAKNTVYLQMNSLKPEDTAVYYCAA
VHH#m9A     VSWSSSTIVYADSVEG  RFTISRDNHQNTVYLQMDSLKPEDTAVYYCAA
VHH#m9E     VSWSGGTIVYADSVKG  RFEISRDNARNTVYLQMDSLKSEDTAVYYCAA
             *   *  *  ** *  ** ****  ***** *** *********

                      CDR3            FR4          Hinge
VHH#m3F     RPYQKYNWA-SASYNV  WGQGTQVTVSS  EPKTPKPQP
VHH#m4B     SP--KYMTAYERSYDF  WGQGTQVTVSS  EPKTPKPQP
VHH#m9A     RPYQKYNWA-SASYNV  WGQGTQVTVSS  ---------
VHH#m9E     RPYQRFNWA-SASYNV  WGRGTQVTVSS  ---------
             *    *  *    **  ** *******
```
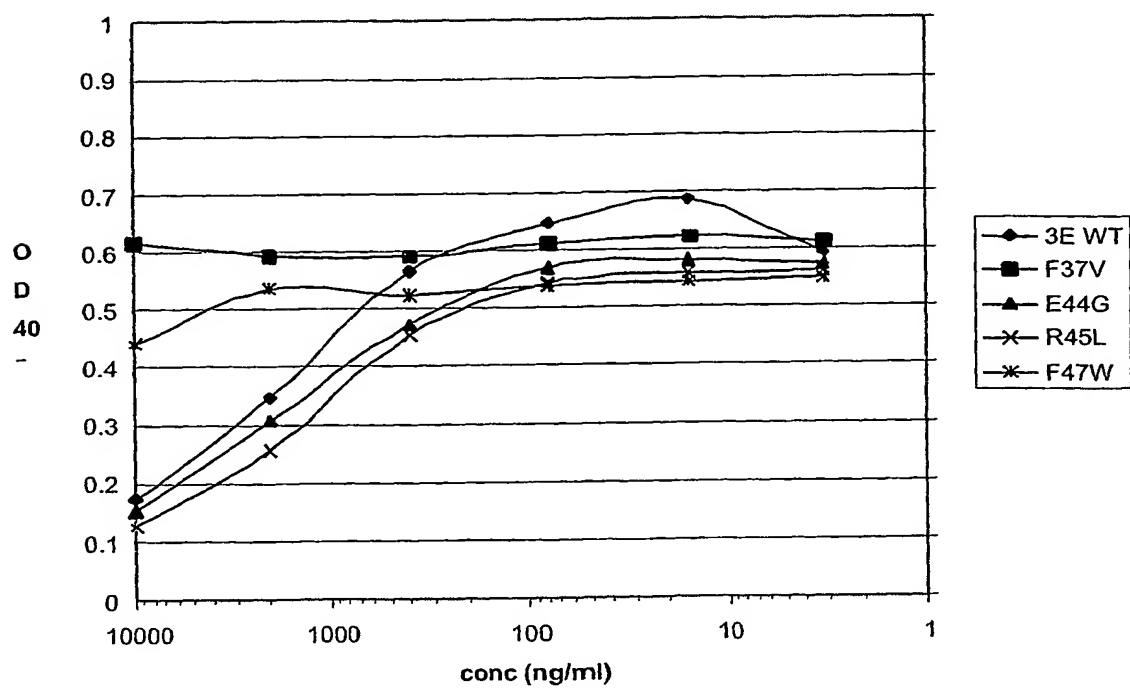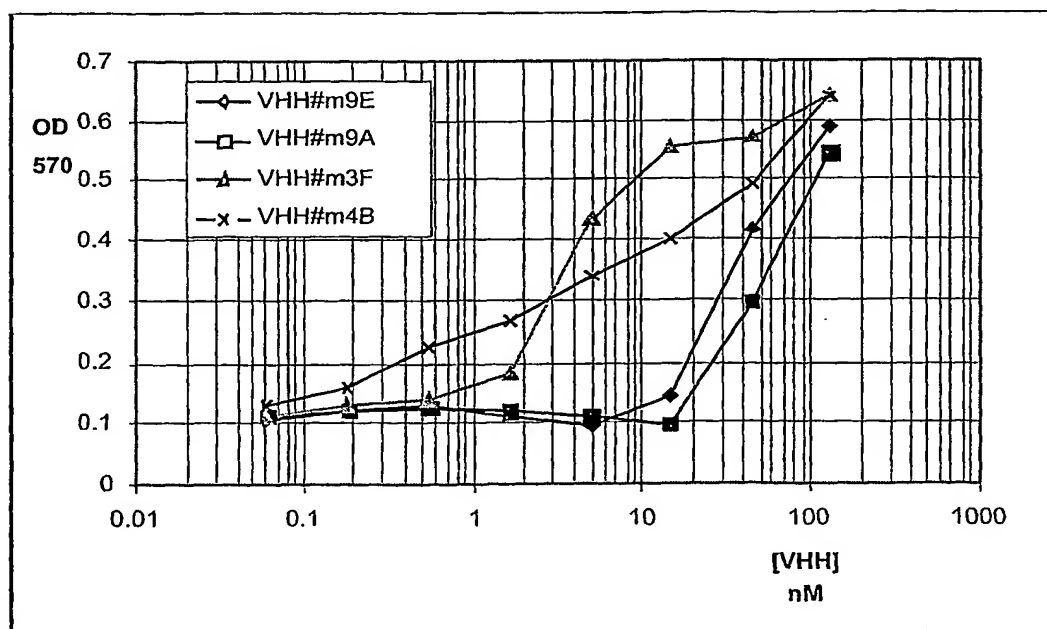
Figure 8

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Figure 9

```
   HindIII
 1 aagcttgcat gcaaattcta tttcaaggag acagtcataa tgaaatacct attgcctacg gcagccgctg gattgttatt
                                                   M  K  Y   L  L  P  T   A  A  A   G  L  L
                                                  <                        pelB-leader

           SfiI    NcoI              NotI                  PstI
81 actcgcggcc cagccggcca tggggcctaa taggcggccg cacaggtgca gctgcaggag tcataatgag ggacccaggt
    L  A  A   Q  P  A   M  G  P -   -  A  A   A  Q  V   Q  L  Q  E   S  -  -    G  T  Q  V
             Leader            ><   VHH#1 >               <                        VHH#2

    BstEII
161 caccgtctcc tcagaacaaa aactcatctc agaagaggat ctgaatgggg ccgcacatca tcatcatcat cattaatgag
     T  V  S   S  E  Q   K  L  I   S  E  E  D   L  N  G   A  A  H   H  H  H  H   H  -  -
             ><         C-MYC                           >           <        His6            >

    EcoRI
241 aattcactgg ccg
```
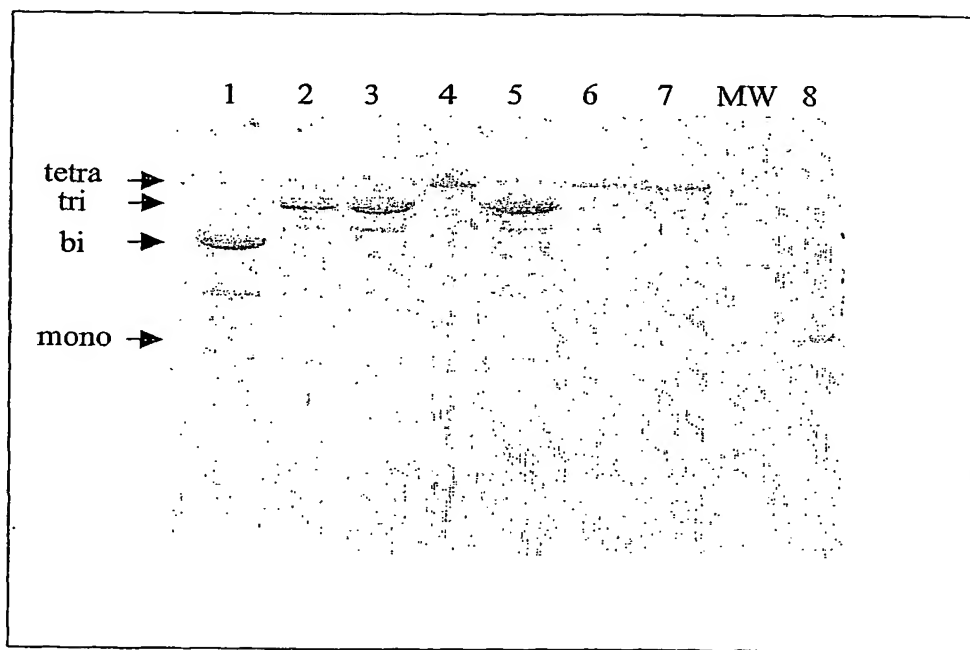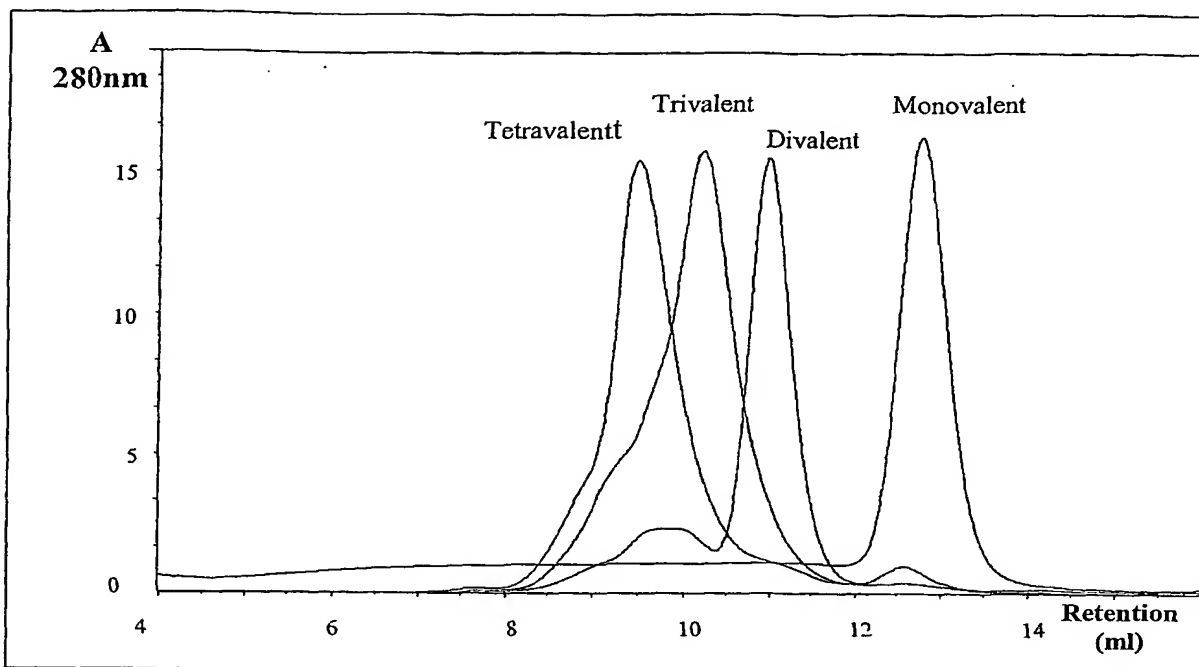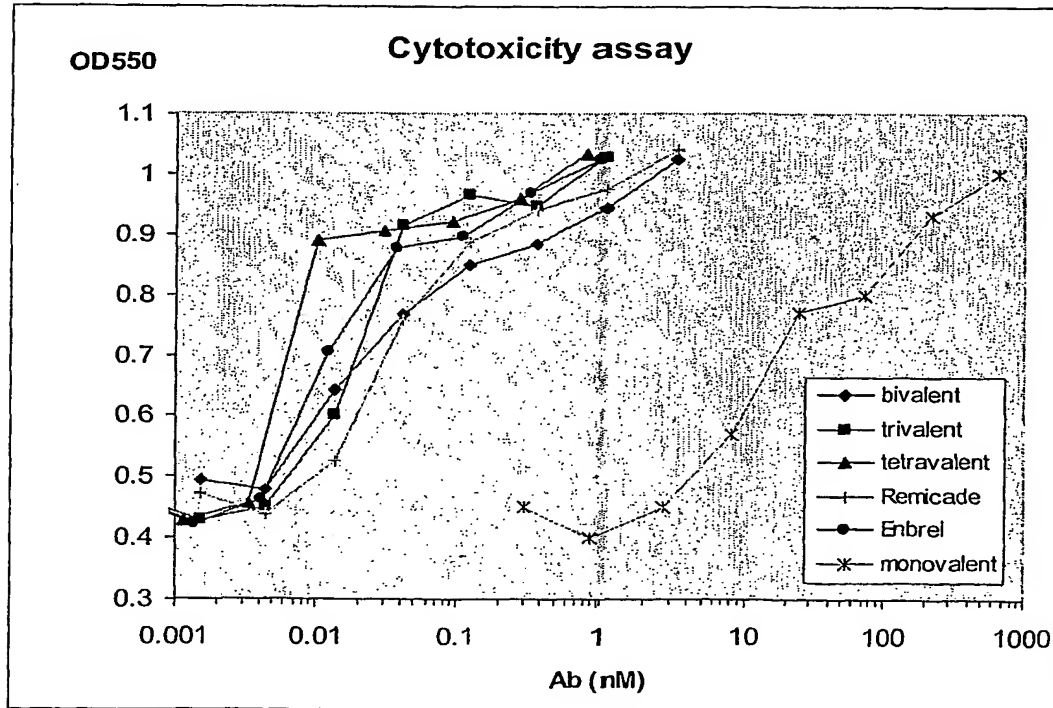
Figure 10

Figure 11

Figure 12

Figure 13
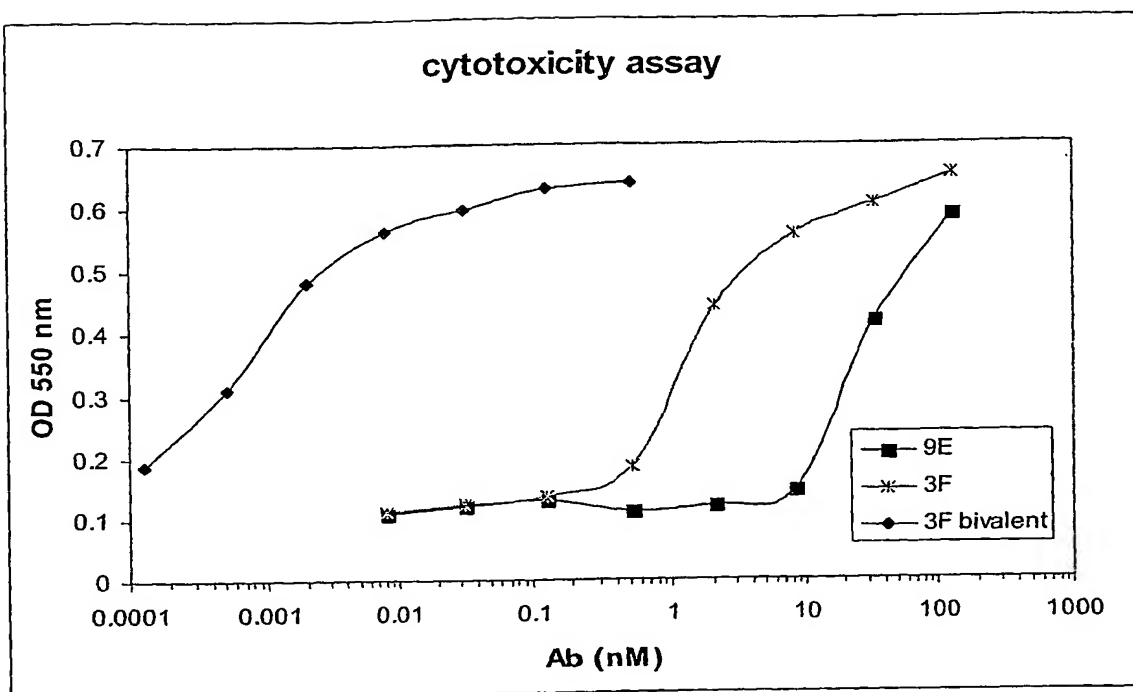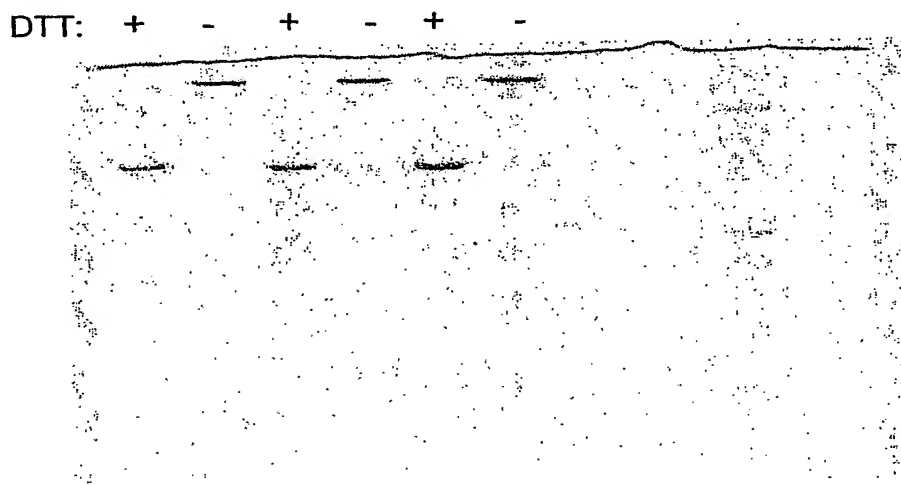
Figure 14

Figure 15
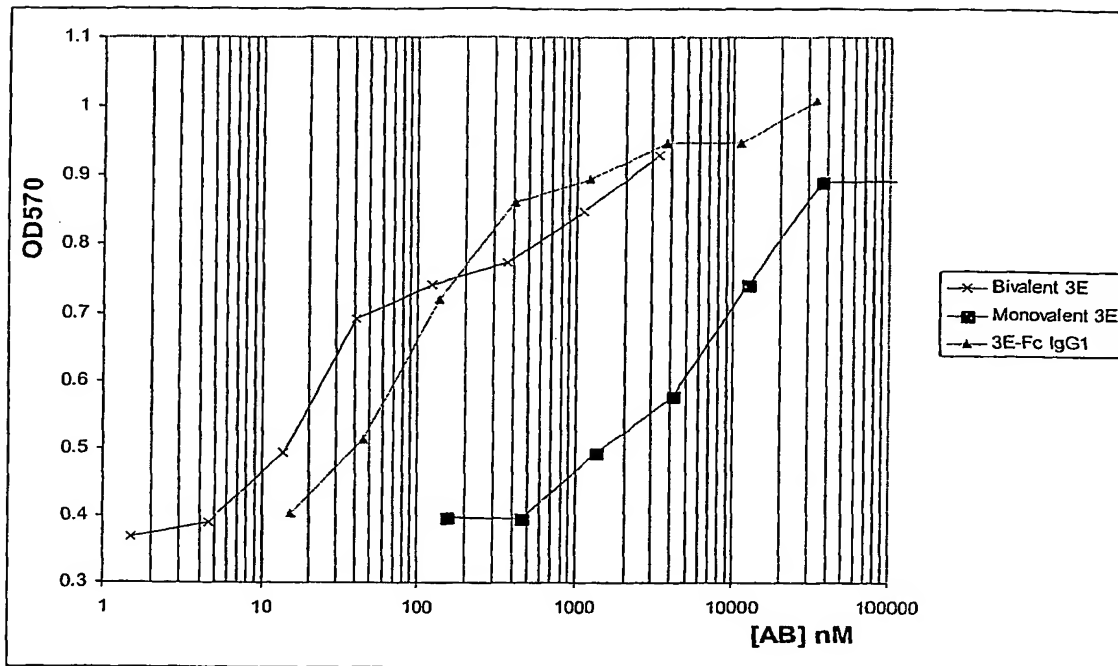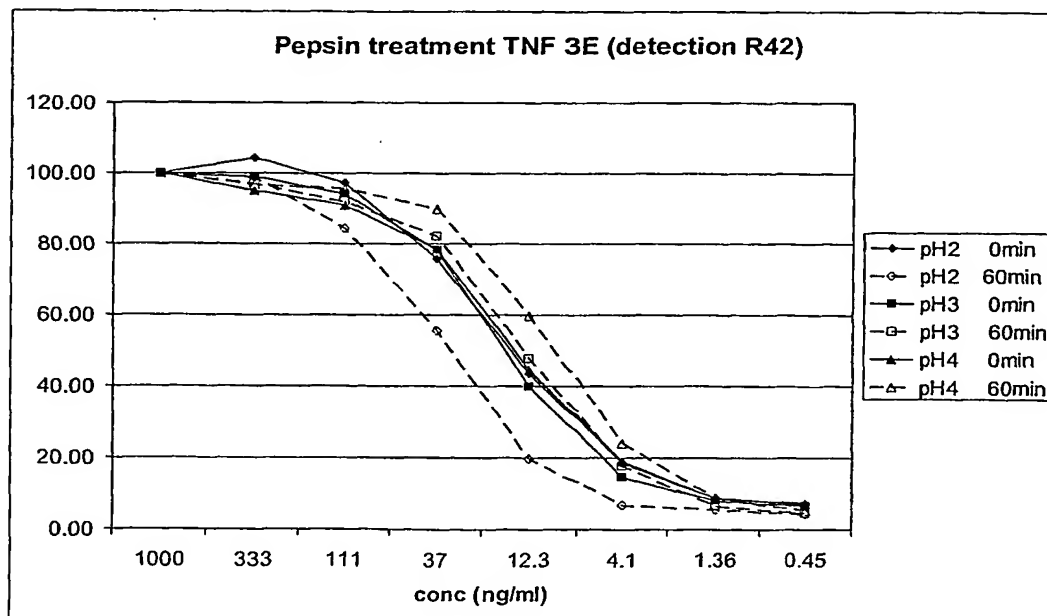
Figure 16

Figure 17

DSS-induced Model of Chronic Colitis

| 7 | 12 | 7 | 12 | 7 | 12 | 7 | 21 | 14 | 7 | 7 |

113 d

7	DSS induction (7 d)
12	Recovery (12 d)
14	Treatment period (14 d)
⇩	Sacrifice of animals
7	Analysis and histopathology

Figure 18